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FIG. 6

PROCESS	WORLD WIDE WEB SITE PAGE
sequence format conversion	<a href="http://dot.imgen-bcm.tmc.edu:9331/seq-util/Options/feadseq.html">http://dot.imgen-bcm.tmc.edu:9331/seq-util/Options/feadseq.html</a>
reverse complementation	<a href="http://dot.imgen-bcm.tmc.edu:9331/seq-util/Options/revcomp.html">http://dot.imgen-bcm.tmc.edu:9331/seq-util/Options/revcomp.html</a>
sequence translation	<a href="http://dot.imgen-bcm.tmc.edu:9331/seq-util/Options/sixframe.html">http://dot.imgen-bcm.tmc.edu:9331/seq-util/Options/sixframe.html</a>
protein information	<a href="http://www.expasy.ch/ttools">http://www.expasy.ch/ttools</a> <i>ExPASy Proteomics Tools</i>
sequence alignments using Clustal W	<a href="http://dot.imgen-bcm.tmc.edu:9331/multi-align/Options/ps/eletetain.html">http://dot.imgen-bcm.tmc.edu:9331/multi-align/Options/ps/eletetain.html</a>
sequence comparisons using BLAST 2.0	<a href="http://www.ncbi.nlm.nih.gov/blast/blast.cgi?jform=0">http://www.ncbi.nlm.nih.gov/blast/blast.cgi?jform=0</a> <i>National Center For Biotechnology</i>
sequence searches using BLAST 2.0	<i>National Center for Biotechnology</i>